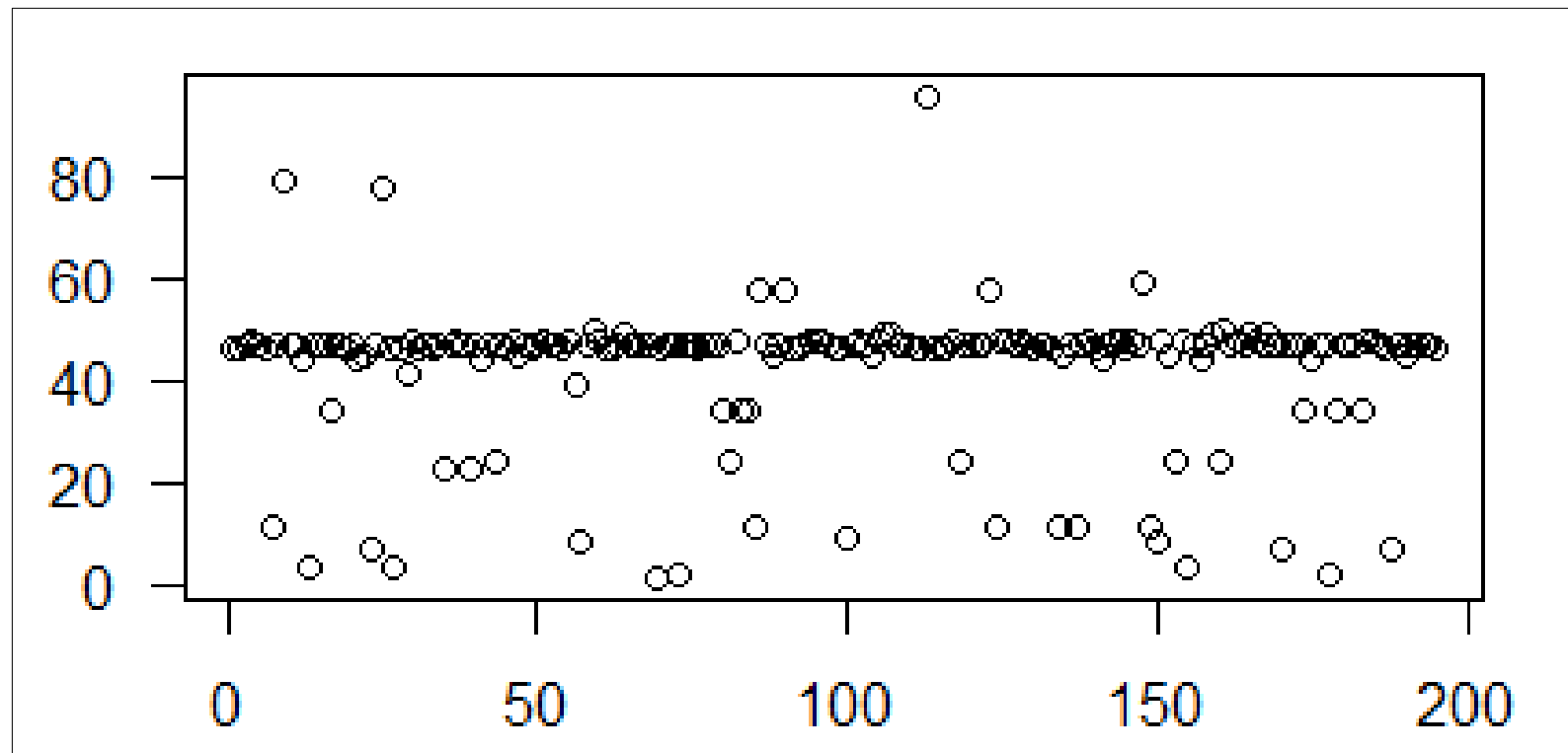
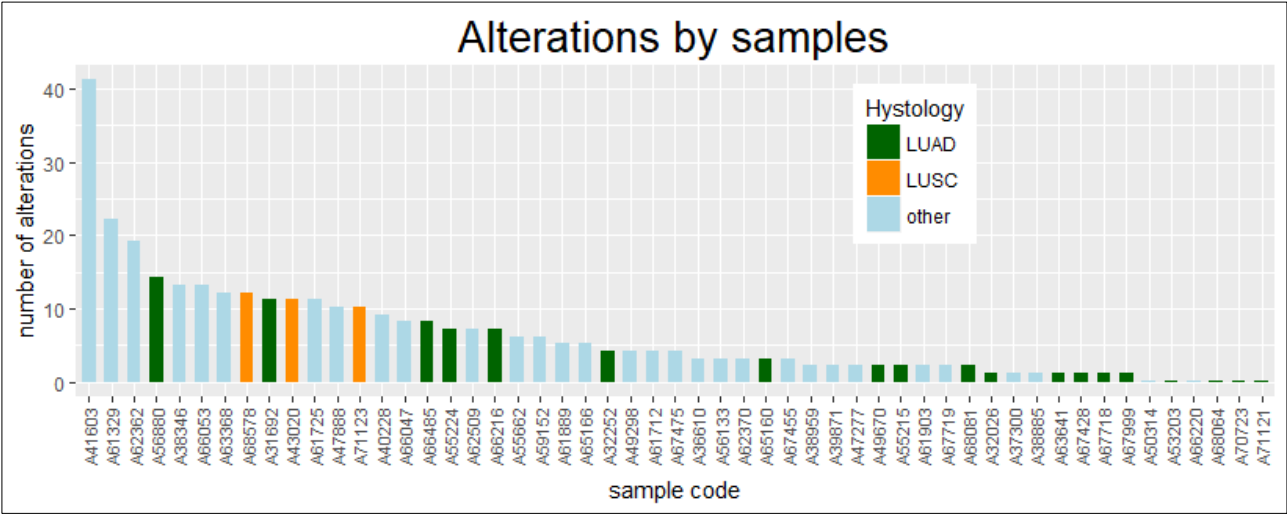


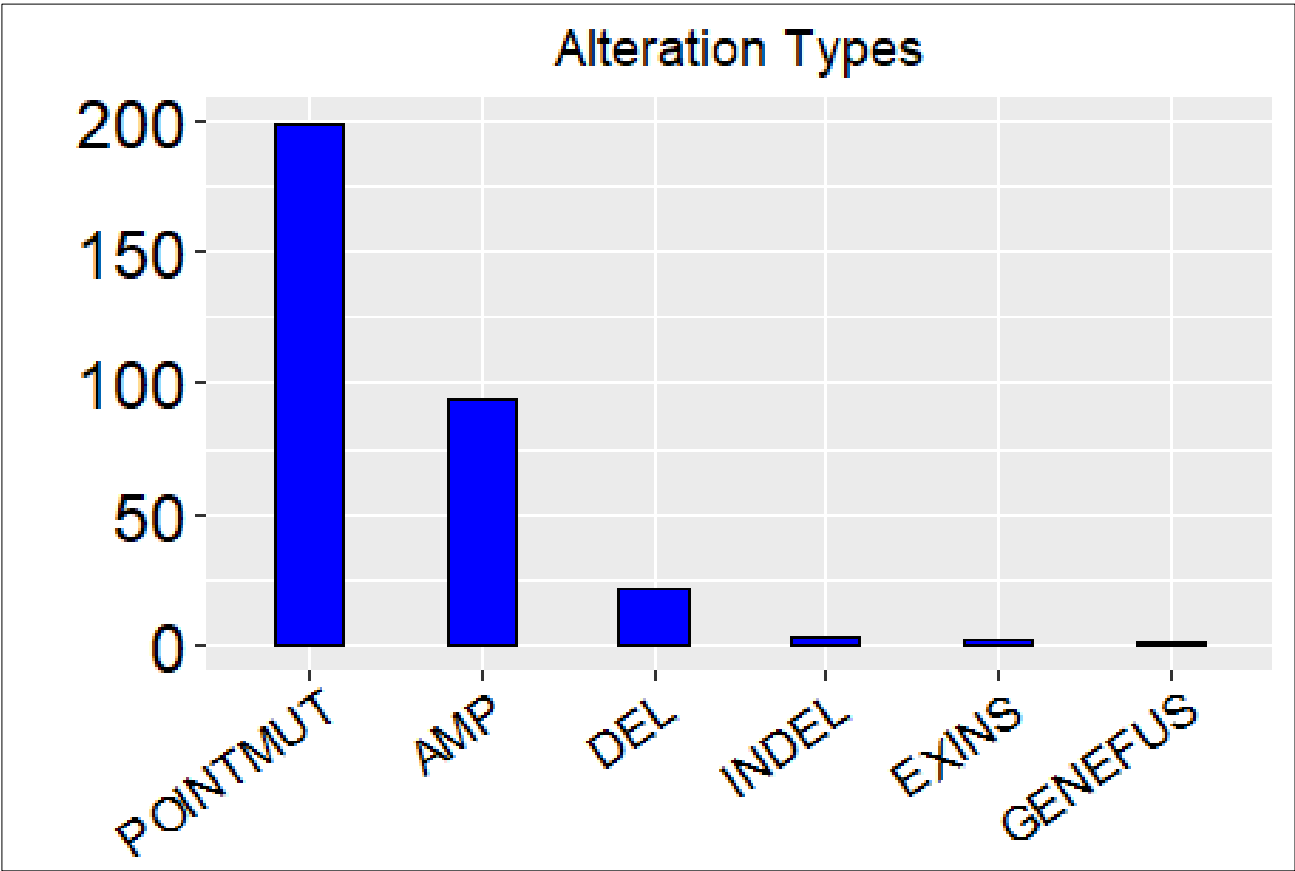
SUPPL. Figure 1: number of genes sequenced in tumor tissue in each patient



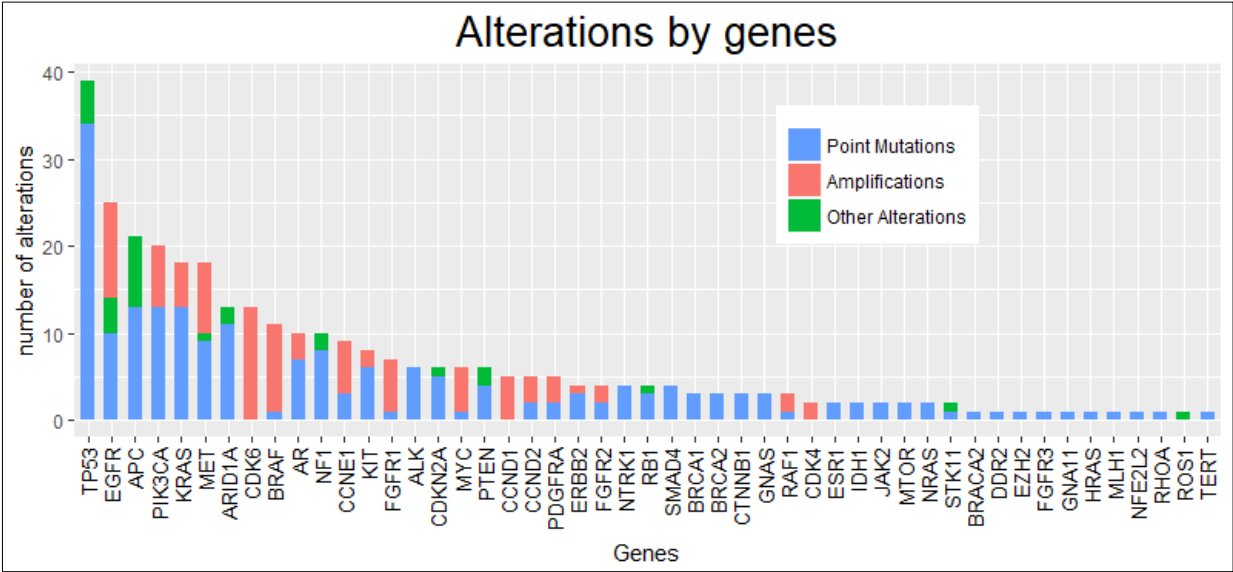
SUPPL. Figure 2: number of alteration by sample in lbNGS patients



SUPPL. Figure 3: types of alterations detected with LB



SUPPL Figure 4: number of alterations by genes in lbNGS samples



SUPPL. Figure 5: frequency of alterations for each gene across all sequenced samples in ttNGS patients (genes sequenced in less than 5 samples and with a alterations frequency < 1% are not represented)

